# nidmviewer Documentation

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This is a command line (to browser) as well as python module for rendering NIDM Results (turtle files) into an interactive viewer. The viewer currently supports parsing and interactive browsing of peak coordinates, viewing associated brain maps, and saving static images to file.

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## CHAPTER 1

## Installation

To install

```
pip install git+git://github.com/vsoch/nidmviewer.git
```

## 1.1 Running Examples

### 1.1.1 Python (server)

```
#!/usr/bin/python
```

### 1.1.2 Command Line

When installing with setup.py, an executable, *nidmviewer* is installed in your bin to view nidm files on the fly:

nidmviewer fsl/nidm.ttl

You can see the basic usage by typing the command:

nidmviewer usage: nidmviewer [-h] [--columns\_to\_remove COLUMNS\_TO\_REMOVE] ttl base nidmviewer: error: too few arguments usage: nidmviewer [-h] [--columns\_to\_remove COLUMNS\_TO\_REMOVE] ttl base command line or server tool to view or compare nidm results. positional arguments: ttl List of comma separated ttl files to parse. base image (standard brain map) to use for the viewer background.

## CHAPTER 2

### nidmviewer

## 2.1 nidmviewer package

### 2.1.1 Subpackages

### 2.1.2 Submodules

### 2.1.3 nidmviewer.browser module

browser.py: part of nidmviewer package Functions to visualize in browser

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nidmviewer.browser.get\_svg\_html (mpl\_figures)

nidmviewer.browser.internal\_view(html\_snippet, tmp\_file)

```
nidmviewer.browser.run_webserver(port=8088, html_page='index.html')
```

```
nidmviewer.browser.view(html_snippet, copy_list, port)
```

```
nidmviewer.browser.write_file(html_snippet, tmp_file)
```

### 2.1.4 nidmviewer.convert module

convert Functions to convert/parse output from nidm sparql queries

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nidmviewer.convert.getjson(nidm\_file, format='n3')

```
nidmviewer.convert.parse_coordinates(coordinates)
```

convert a list of xyz strings in format '[x,y,z]' to separate variables This is what we get from the sparql query Parameters ======== coordinates: list

a list of xyz coordinate strings, each a list in a string '[x,y,z]'

coordinate\_df: pandas dataframe columns are x,y,z

```
nidmviewer.convert.prettyjson(nidm_file, format='n3')
```

### 2.1.5 nidmviewer.scripts module

script.py: part of nidmviewer package Functions to visualize in browser

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nidmviewer.scripts.main()

#### 2.1.6 nidmviewer.sparql module

sparql.py: part of the nidmviewer package Sparql queries

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nidmviewer.sparql.do\_query (ttl\_file, query, rdf\_format='turtle', serialize\_format='csv', output\_df=True)

nidmviewer.sparql.get\_coordinates(ttl\_file)

nidmviewer.sparql.get\_coordinates\_and\_maps(ttl\_file)

### 2.1.7 nidmviewer.templates module

templates.py: part of the nidmviewer package Functions to work with html templates

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nidmviewer.templates.add\_javascript\_function(function\_code, template)

nidmviewer.templates.add\_string(tag, substitution, template)
nidmviewer.templates.get\_image(image\_name)
nidmviewer.templates.get\_template(html\_name)
nidmviewer.templates.read\_template(html\_name)
nidmviewer.templates.remove\_resources(html\_snippet, script\_names)
nidmviewer.templates.save\_template(html\_snippet, output\_file)

### 2.1.8 nidmviewer.utils module

utils.py: part of the nidmviewer package Functions to work with html templates

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```
nidmviewer.utils.download_file(src, dest)
```

```
nidmviewer.utils.get_extension(path)
```

nidmviewer.utils.get\_images (peaks, location\_key)
get\_images returns unique images for a location key from a peaks table

nidmviewer.utils.get\_name(path)

nidmviewer.utils.get\_package\_dir()

nidmviewer.utils.get\_random\_name(length=6, chars='ABCDEFGHIJKLMNOPQRSTUVWXYZ0123456789')

```
nidmviewer.utils.get_standard_brain(load=True)
```

nidmviewer.utils.is\_empty(nii\_file)

nidmviewer.utils.make\_dir(directory)

nidmviewer.utils.make\_png\_paths (nifti\_files)

nidmviewer.utils.make\_tmp\_folder(\*args, \*\*kwds)

nidmviewer.utils.read\_file\_lines(file\_name)

```
nidmviewer.utils.strip_url(url, encode=True)
```

nidmviewer.utils.unwrap\_list\_unique(list\_of\_lists)

```
nidmviewer.utils.unzip(source, dest_dir)
```

#### 2.1.9 nidmviewer.viewer module

viewer.py: part of the nidmviewer package

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nidmviewer.viewer.check\_inputs(ttl\_files)

full paths to ttl files

list of ttl files

one or more turtle files to add to the viewer. Images in the files should be available at the specified URL.

- **retrieve: boolean** If set to False, the images are assumed to be on the same server, and will be served with the given URL. If retrieve is set to True, the images will be retrieved first and stored in a temporary directory.
- **base\_image: str** The base image to use for the viewer. Not specifying a base\_image will yield a black back-ground.
- view\_in\_browser: boolean open a temporary web browser (to run locally). If True, images will be copied to a temp folder. If False, image\_paths must be relative to web server. File names should be unique.
- columns\_to\_remove: additional columns to remove. If none, default columns of "coordinate\_id" "statmap\_type" and "exc\_set" will be removed.
- **port: int** port to serve nidmviewer, if view\_in\_browser==True
- **remove\_scripts: list** one or more script or button tags to remove from the template. Options include JQUERY BOOTSTRAPJS BOOTSTRAPCSS PAPAYACSS PAPAYAJS NIDMSELECTBUTTON

button\_text: str Text string for the button to select a brain image. Default is "BRAIN"

check\_empty: boolean - check for empty images or not. Will result in error if nidm paths are URLS.

nidmviewer.viewer.generate\_temp (peaks, location\_key)

generate a lookup of temporary files Parameters ======= peaks: dict

data structure from get\_coordinates\_and\_peaks

location\_key: str key in peaks data structure for file paths

- **peaks: dict** (key is ttl file, equivalent to peaks, but old location\_key path is replaced with path to temporary directory
- **copy\_list: dict** keys are current paths, values are temporary file names corresponding to fullpath in new\_nifti\_files[ttl\_file] dictionary. This is used to copy images into the temporary directory with the correct names.

nidmviewer.viewer.get\_column\_names(peaks)

nidmviewer.viewer.parse\_nidm(ttl\_files)

Extract brainmaps and coordinates from ttl files Parameters: ======== ttl\_files: list

list of full paths to ttl files

- **peaks: dict** dict of pandas data frames, one for each ttl\_file, with columns coordinate, z, peak\_name, pvalue\_uncorrected
- **maps: dict** dict of pandas data frames, one for each ttl\_file, with columns filename and location for all brain maps specified in ttl.
- nidmviewer.viewer.remove\_columns (columns, columns\_to\_remove)
- - dictionary (key, is ttl\_file, and value, is dictionary of {filename:fullpath} for all brainmaps extracted from the ttl files
  - retrieve: boolean if True, will download brainmaps to temporary folder first. If false, encodes path in utf-8 for rendering in javascript

location\_key: str key to look up file name in peaks dictionary

data frame to convert

- orient: str orientation to convert with (default is "records")
- strip\_columns: boolean if true, will return df.to\_dict(orient=orient).values() default is False

strings: boolean True will convert all columns to strings.

### 2.1.10 Module contents

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